

## RESULT 12

AAV24010

ID AAV24010 standard; DNA; 2157 BP.

AC AAV24010;

DT 06-AUG-1998 (first entry)

DE Human FK506 binding protein, FKBP52, coding sequence.

KW FK506 binding protein; FKBP52; human; immune response regulator;  
KW immunosuppressant; steroid hormone receptor transformation; ds.

XX Homo sapiens.

XX Key Location/Qualifiers  
FH 100..1479FT CDS  
FT /\*tag= a  
FT /product= "FKBP52"

XX US5763590-A.

XX 09-JUN-1998.

XX 09-NOV-1994; 94US-00336618.

XX 11-OCT-1991; 91US-00777752.

PR 16-OCT-1992; 92US-00963325.

PR 29-MAR-1994; 94US-00218989.

XX (VERT-) VERTEX PHARM INC.

XX Peattie DA, Livingston DJ, Harding MW;

WPI; 1998-347419/30.

P-PSDB; AAW54038.

XX DNA sequence encoding human FK506-binding protein - and recombinant DNA  
XX molecule containing it.

XX Claim 1; Fig 3; 28pp; English.

XX This sequence encodes 52 kD human FK506 binding protein, referred to as  
XX FKBP52, of the invention. The FKBP52 protein plays a key role in  
XX regulating immune responses. FKBP52 may be useful for mediating steroid  
XX hormone receptor transformation. The DNA may be used to screen for new  
XX immunosuppressants, and in assays for metabolites in samples from  
XX individuals taking immunosuppressants. The DNA may also be used in assays  
XX for identifying natural intracellular rapamycin-like or FK506 like  
XX substances, and in assays for identifying natural intracellular  
XX substrates that are potential targets for other immunosuppressants

SQ Sequence 2157 BP; 531 A; 551 C; 606 G; 469 T; 0 U; 0 Other;

102665

Query Match 4.5%; Score 56.8; DB 2; Length 2157;  
 Best Local Similarity 47.5%; Pred. No. 1.1e-05;  
 Matches 280; Conservative 0; Mismatches 292; Indels 18; Gaps 3;

QY 471 GGTATTAGCCATCGGTGTTCTAGCATGAAGTCTGGTGAACGTGCGCTGTGTGATGTTGGC 530  
 DB |||||  
 QY 706 GGTCTGGAGAGGCCATTTCAGCGCATGGAGAAAGAGACATTCATCGTGTACCTCAAG 765  
 DB |||||  
 QY 531 TGGGAATTAGCTTATGGGAAGAGAACTTTCTTTTCCCAATGTTCCACCTATGGCA 590  
 DB |||||  
 QY 766 CCCAGCTATGCTTTTGGCAGTGTGGGAAGGAAAGTT---CCAAATCCCAAAATGCT 822  
 DB |||||  
 QY 591 GACTTGTATTATGAGTGAAGTTATTTGGTTTGTATGAACAAGAGAGGAAAGCTGCG 650  
 DB |||||  
 QY 823 GAGCTGAATATGAATTAACCTCAAGAGTTTGAAGGCGCAAGGAG-----TCT 873  
 DB |||||  
 QY 651 AGTGATATGACTGTAGAGAAAGGATTTGGTGCAGCAGACAGAAAGAAATGATGGGAAT 710  
 DB |||||  
 QY 874 TGGGAGATGAATTCAGAGAGAGCTGGNACAGAGCACCATAGTGAAGAGCGGGGCACT 933  
 DB |||||  
 QY 711 TCTCTTTTAAAGGAGAGAACTGGAGGAAGCCATGCAACAGTATGAATGGCCATAGCA 770  
 DB |||||  
 QY 934 GTGTACTTTCAGGAAGGTAATAACAAGCAAGCTTTTACTACAGTATAAGAGATCGTGTCT 993  
 DB |||||  
 QY 771 TACATGGGGGACGATTTTATGTTTTCAGCTGTATGGGAAGTACCAGGATATGGCTTTAGCA 830  
 DB |||||  
 QY 994 TGGCTGGAAATATGATCTAGTGT-----TCCAAATGAGGAAGCACAGAAAGCACAGGCC 1047  
 DB |||||  
 QY 831 GTTAAAAACCCATGCCATCTTAACATAGCAGCTTGCCTCATCAAACTTAAACGATACGAT 890  
 DB |||||  
 QY 1048 CTTCGACTGGCCTCTCACCTCAACCTGGCCATGTGTCTCTGAACTACAGGCTTCTCT 1107  
 DB |||||  
 QY 891 GAAGCAATTTGGTCACTGCAACATTTGTGTGAAGAGAAAGAGAAACCCAAAGCACTG 950  
 DB |||||  
 QY 1108 GCTGCCATTTGAAGAGCTGTAAAGAGCCCTAGAACTGGACAGCAACAACGAGAGGGCCTC 1167  
 DB |||||  
 QY 951 TTCAAGAGAGGAAAGCAAGGAGCAGCTAGGACAGATGGACTCAGCAGCTGATGATTTTC 1010  
 DB |||||  
 QY 1168 TTCCCGGGGAGAGGCCCACTGGCGGTGAATGATTTGAACCTGGCAGCGGCTGATTTTC 1227  
 DB |||||  
 QY 1011 CGAAAGGCAAAAGTATGCTCTCTGACGACAGAGGCGGATTTAGAGAGAGCT 1060  
 DB |||||  
 QY 1228 CAGAAGGTCTCTGAGCTCTACCCCAACAAAGCGCGCCAGACCAGCT 1277  
 DB |||||

XX  
 mn wpt...2002-740862/80.